

BRIEF DESCRIPTION OF THE FIGURES

Figures 1A-1C

5 Nucleotide sequence including sequence encoding a
human SNORF68 receptor (SEQ ID NO: 1). Putative open
reading frames including the shortest open reading
frame are indicated by underlining two start (ATG)
codons (at positions 26-28 and 62-64) and the stop
codon (at positions 1508-1510). In addition, partial
10 5' and 3' untranslated sequences are shown.

Figures 2A-2C

15 Deduced amino acid sequence (SEQ ID NO: 2) of the
human SNORF68 receptor encoded by the longest open
reading frame indicated in the nucleotide sequence
shown in Figures 1A-1C (SEQ ID NO: 1). The seven
putative transmembrane (TM) regions are underlined.